RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/016,590
Source: 1Fw16
Date Processed by STIC: 3/10/05

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IFW16

RAW SEQUENCE LISTING DATE: 03/10/2005
PATENT APPLICATION: **US/09/016,159D** TIME: 10:17:03

Input Set : **D:\completeseq2.txt**
Output Set: **N:\CRF4\03102005\I016159D.raw**

```
 3 <110> APPLICANT: Lee, Jong Y.
 5 <120> TITLE OF INVENTION: PURIFIED HUMAN ERYTHROPOIETIN RECEPTOR PROTEIN FRAGMENT AND
 6       ANTIBODIES DERIVED THEREFROM
 8 <130> FILE REFERENCE: 106.001US2
10 <140> CURRENT APPLICATION NUMBER: US 09/016,159D
11 <141> CURRENT FILING DATE: 1998-01-30
13 <150> PRIOR APPLICATION NUMBER: US 08/876,227
14 <151> PRIOR FILING DATE: 1997-06-16
16 <160> NUMBER OF SEQ ID NOS: 7
18 <170> SOFTWARE: PatentIn version 3.3
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 23
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial
25 <220> FEATURE:
26 <223> OTHER INFORMATION: BamH1 linker at 5' end followed by sequence for amino acids
25
27       through 29 of full length EpoR protein. Forward primer for SEQ
28       ID NO:2.
30 <400> SEQUENCE: 1
31 ttggatccgc gcccccgcct aac                                          23
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 22
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial
39 <220> FEATURE:
40 <223> OTHER INFORMATION: EcoR1 linker followed by sequence complementary to coding
41       sequence for amino acids 226 through 222 of full length human
42       EpoR protein. Reverse primer for SEQ ID NO:1.
44 <400> SEQUENCE: 2
45 tgaattcggg gtccaggtcg ct                                           22
48 <210> SEQ ID NO: 3
49 <211> LENGTH: 18
50 <212> TYPE: DNA
51 <213> ORGANISM: Homo sapiens
53 <300> PUBLICATION INFORMATION:
54 <301> AUTHORs: Smith, D.B. et al.
55 <302> TITLE: Single-step purification of polypeptides expressed in Escherichia
56       coli as fusions with glutathione-S-transferase
57 <303> JOURNAL: Gene
58 <304> VOLUME: 67
59 <306> PAGES: 31-40
60 <307> DATE: 1998
62 <300> PUBLICATION INFORMATION:
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<301> AUTHORs: Smith, D.B. et al.
<302> TITLE: Single-step purification of polypeptides expressed in Escherichia
coli as fusions with glutathione-S-transferase
<303> JOURNAL: Genes and Development
<304> VOLUME: 67
<306> PAGES: 31-40
<307> DATE: 1998
<400> SEQUENCE: 3
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<210> SEQ ID NO: 4
<211> LENGTH: 1527
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens
<300> PUBLICATION INFORMATION:
<301> AUTHORs: Jones, S.S. et al.
<302> TITLE: Human Erythropoietin Receptor: Cloning, expression, and
biological characterization
<303> JOURNAL: Blood
<304> VOLUME: 76
<305> ISSUE: 1
<306> PAGES: 31-35
<307> DATE: 1990-07-01
<400> SEQUENCE: 4
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ccccccagct atgtggcttg ctcttag 1527

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Input Set : **D:\completeseq2.txt**
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<210> SEQ ID NO: 5
<211> LENGTH: 508
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<300> PUBLICATION INFORMATION:
<301> AUTHORs: Jones, S.S. et al.
<302> TITLE: Human Erythropoietin Receptor: Cloning, expression, and biological characterization
<303> JOURNAL: Blood
<304> VOLUME: 76
<305> ISSUE: 1
<306> PAGES: 31-35
<307> DATE: 1990-07-01
<400> SEQUENCE: 5

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1               5                  10                  15
Leu Leu Leu Ala Gly Ala Ala Trp Ala Pro Pro Pro Asn Leu Pro Asp
            20                  25                  30
Pro Lys Phe Glu Ser Lys Ala Ala Leu Leu Ala Ala Arg Gly Pro Glu
        35                  40                  45
Glu Leu Leu Cys Phe Thr Glu Arg Leu Glu Asp Leu Val Cys Phe Trp
    50                  55                  60
Glu Glu Ala Ala Ser Ala Gly Val Gly Pro Gly Asn Tyr Ser Phe Ser
65                  70                  75                  80
Tyr Gln Leu Glu Asp Glu Pro Trp Lys Leu Cys Arg Leu His Gln Ala
                85                  90                  95
Pro Thr Ala Arg Gly Ala Val Arg Phe Trp Cys Ser Leu Pro Thr Ala
            100                 105                 110
Asp Thr Ser Ser Phe Val Pro Leu Glu Leu Arg Val Thr Ala Ala Ser
        115                 120                 125
Gly Ala Pro Arg Tyr His Arg Val Ile His Ile Asn Glu Val Val Leu
    130                 135                 140
Leu Asp Ala Pro Val Gly Leu Val Ala Arg Leu Ala Asp Glu Ser Gly
145                 150                 155                 160
His Val Val Leu Arg Trp Leu Pro Pro Pro Glu Thr Pro Met Thr Ser
                165                 170                 175
His Ile Arg Tyr Glu Val Asp Val Ser Ala Gly Asn Gly Ala Gly Ser
            180                 185                 190
Val Gln Arg Val Glu Ile Leu Glu Gly Arg Thr Glu Cys Val Leu Ser
        195                 200                 205
Asn Leu Arg Gly Arg Thr Arg Tyr Thr Phe Ala Val Arg Ala Arg Met
    210                 215                 220
Ala Glu Pro Ser Phe Gly Gly Phe Trp Ser Ala Trp Ser Glu Pro Val
225                 230                 235                 240
Ser Leu Leu Thr Pro Ser Asp Leu Asp Pro Leu Ile Leu Thr Leu Ser
                245                 250                 255
Leu Ile Leu Val Val Ile Leu Val Leu Leu Thr Val Leu Ala Leu Leu
            260                 265                 270
Ser His Arg Arg Ala Leu Lys Gln Lys Ile Trp Pro Gly Ile Pro Ser
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233 Pro Glu Ser Glu Phe Glu Gly Leu Phe Thr Thr His Lys Gly Asn Phe
234     290                 295                 300
237 Gln Leu Trp Leu Tyr Gln Asn Asp Gly Cys Leu Trp Trp Ser Pro Cys
238 305                 310                 315                 320
241 Thr Pro Phe Thr Glu Asp Pro Pro Ala Ser Leu Glu Val Leu Ser Glu
242                 325                 330                 335
245 Arg Cys Trp Gly Thr Met Gln Ala Val Glu Pro Gly Thr Asp Asp Glu
246             340                 345                 350
249 Gly Pro Leu Leu Glu Pro Val Gly Ser Glu His Ala Gln Asp Thr Tyr
250         355                 360                 365
253 Leu Val Leu Asp Lys Trp Leu Leu Pro Arg Asn Pro Pro Ser Glu Asp
254     370                 375                 380
257 Leu Pro Gly Pro Gly Gly Ser Val Asp Ile Val Ala Met Asp Glu Gly
258 385                 390                 395                 400
261 Ser Glu Ala Ser Ser Cys Ser Ser Ala Leu Ala Ser Lys Pro Ser Pro
262                 405                 410                 415
265 Glu Gly Ala Ser Ala Ala Ser Phe Glu Tyr Thr Ile Leu Asp Pro Ser
266             420                 425                 430
269 Ser Gln Leu Leu Arg Pro Trp Thr Leu Cys Pro Glu Leu Pro Pro Thr
270         435                 440                 445
273 Pro Pro His Leu Lys Tyr Leu Tyr Leu Val Val Ser Asp Ser Gly Ile
274     450                 455                 460
277 Ser Thr Asp Tyr Ser Ser Gly Asp Ser Gln Gly Ala Gln Gly Gly Leu
278 465                 470                 475                 480
281 Ser Asp Gly Pro Tyr Ser Asn Pro Tyr Glu Asn Ser Leu Ile Pro Ala
282                 485                 490                 495
285 Ala Glu Pro Leu Pro Pro Ser Tyr Val Ala Cys Ser
286             500                 505
289 <210> SEQ ID NO: 6
290 <211> LENGTH: 1527
291 <212> TYPE: DNA
292 <213> ORGANISM: Homo sapiens
294 <300> PUBLICATION INFORMATION:
295 <301> AUTHORs: Winkelman, J.C. et al.
296 <302> TITLE: The gene for the human erythropoietin receptor: analysis of the
297       coding sequence and assignment to chromosome 19p
298 <303> JOURNAL: Blood
299 <304> VOLUME: 76
300 <305> ISSUE: 1
301 <306> PAGES: 24-30
302 <307> DATE: 1990-07-01
304 <400> SEQUENCE: 6
305 atggaccacc tcggggcgtc cctctggccc caggtcggct ccctttgtct cctgctcgct     60
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309 ttgctggcgg cccgggggcc cgaagagctt ctgtgcttca ccgagcggtt ggaggacttg    180
311 gtgtgtttct gggaggaagc ggcgagcgct ggggtgggcc cgggcaacta cagcttctcc    240
313 taccagctcg aggatgagcc atggaagctg tgtcgcctgc accaggctcc cacggctcgt    300
315 ggtcgggtgc gcttctggtg ttcgctgcct acagccgaca cgtcgagctt cgtgccccta    360
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317 gagttgcgcg tcacagcagc ctccggcgct ccgcgatatc accgtgtcat ccacatcaat 420
319 gaagtagtgc tcctagacgc ccccgtgggg ctggtggcgc ggttggctga cgagagcggc 480
321 cacgtagtgt tgcgctggct cccgccgcct gagacaccca tgacgtctca catccgctac 540
323 gaggtggacg tctcggccgg caaccggcca gggagcgtac agagggtgga gatcctggag 600
325 ggccgcaccg agtgtgtgct gagcaacctg cggggccgga cgcgctacac cttcgccgtc 660
327 cgcgcgcgta tggctgagcc gagcttcggc ggcttctgga gcgcctggtc ggagcctgtg 720
329 tcgctgctgg agcctagcga cctggacccc ctcatcctga cgctctccct catcctcgtg 780
331 gtcatcctgg tgctgctgac cgtgctcgcg ctgctctccc accgccgggc tctgaagcag 840
333 aagatctggc ctggcatccc gagcccagag agcgagtttg aaggcctctt caccacccac 900
335 aagggtaact tccagctgtg gctgtaccag aatgatggct gcctgtggtg gagcccctgc 960
337 acccccttca cggaggaccc acctgcttcc ctggaagtcc tctcagagcg ctgctggggg 1020
339 acgatgcagg cagtggagcc ggggacagat gatgagggcc ccctgctgga gccagtgggc 1080
341 agtgagcatg cccaggatac ctatctggtg ctggacaaat ggttgctgcc ccggaacccg 1140
343 cccagtgagg acctcccagg gcctggtggc agtgtggaca tagtggccat ggatgaaggc 1200
345 tcagaagcat cctcctgctc atctgctttg gcctcgaagc ccagcccaga gggagcctct 1260
347 gctgccagct ttgagtacac tatcctggac cccagctccc agctcttgcg tccatggaca 1320
349 ctgtgccctg agctgccccc taccccaccc cacctaaagt acctgtacct tgtggtatct 1380
351 gactctggca tctcaactga ctacagctca ggggactccc agggagccca agggggctta 1440
353 tccgatgggc cctactccaa cccttatgag aacagcctta tcccagccgc tgagcctctg 1500
355 ccccccagct atgtggcttg ctcttag 1527
358 <210> SEQ ID NO: 7
359 <211> LENGTH: 508
360 <212> TYPE: PRT
361 <213> ORGANISM: Homo sapiens
363 <300> PUBLICATION INFORMATION:
364 <301> AUTHORs: Winkelmann, J.C. et al.
365 <302> TITLE: The Gene for the Human Erythropoietin Receptor: Analysis of the
366 coding sequence and assignment to chromosome 19p
367 <303> JOURNAL: Blood
368 <304> VOLUME: 76
369 <305> ISSUE: 1
370 <306> PAGES: 24-30
371 <307> DATE: 1990-07-01
373 <400> SEQUENCE: 7
375 Met Asp His Leu Gly Ala Ser Leu Trp Pro Gln Val Gly Ser Leu Cys
376 1 5 10 15
379 Leu Leu Leu Ala Gly Ala Ala Trp Ala Pro Pro Pro Asn Leu Pro Asp
380 20 25 30
383 Pro Lys Phe Glu Ser Lys Ala Ala Leu Leu Ala Ala Arg Gly Pro Glu
384 35 40 45
387 Glu Leu Leu Cys Phe Thr Glu Arg Leu Glu Asp Leu Val Cys Phe Trp
388 50 55 60
391 Glu Glu Ala Ala Ser Ala Gly Val Gly Pro Gly Asn Tyr Ser Phe Ser
392 65 70 75 80
395 Tyr Gln Leu Glu Asp Glu Pro Trp Lys Leu Cys Arg Leu His Gln Ala
396 85 90 95
399 Pro Thr Ala Arg Gly Arg Val Arg Phe Trp Cys Ser Leu Pro Thr Ala
400 100 105 110
403 Asp Thr Ser Ser Phe Val Pro Leu Glu Leu Arg Val Thr Ala Ala Ser

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/10/2005
PATENT APPLICATION: **US/09/016,159D** TIME: 10:17:04

Input Set : **D:\completeseq2.txt**
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Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2

VERIFICATION SUMMARY DATE: 03/10/2005
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